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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/711,022

DATE: 11/29/2000
 TIME: 09:12:34

ENTERED

Input Set : A:\V1397028.txt
 Output Set: N:\CRF3\11292000\I711022.raw

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4 <110> APPLICANT: MARTHA K. NEWELL
7 <120> TITLE OF INVENTION: METHODS AND PRODUCTS RELATED TO
8   METABOLIC INTERACTIONS IN DISEASE
11 <130> FILE REFERENCE: V0139/7028
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/711,022
C--> 13 <141> CURRENT FILING DATE: 2000-11-09
13 <150> PRIOR APPLICATION NUMBER: U.S. 60/082,250
14 <151> PRIOR FILING DATE: 1998-04-17
16 <150> PRIOR APPLICATION NUMBER: U.S. 60/094,519
17 <151> PRIOR FILING DATE: 1998-07-29
19 <150> PRIOR APPLICATION NUMBER: U.S. 60/101,580
20 <151> PRIOR FILING DATE: 1998-09-24
22 <160> NUMBER OF SEQ ID NOS: 13
24 <170> SOFTWARE: FastSEQ for Windows Version 3.0
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 1491
28 <212> TYPE: DNA
29 <213> ORGANISM: Homo Sapiens
31 <400> SEQUENCE: 1
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33 ggagtctttac cctgaaatca aaggatttaa agaaaaagtg gaatttttct tcagcaagct      120
34 gtgaactata atcccauacc ttggagagcc caggaaacac ctccaatctc tgtgtgtttt      180
35 gtaaacatca ctggagggtc ttctacgtga gcaattggat tgtcatcagc cctgectgtt      240
36 ttgcacctgg gaagtgcctt ggtcttactt ggggccaaat tgttggtctt cacttttgac      300
37 cctaaqcatc tgaagccatg ggccacacac ggaggcaggg aacatcacca tccaagtgtc      360
38 cataacctcaa ttcttttcag ctcttgggtc tggctgggtc ttctacttc tgttcagggt      420
39 ttatccacgt gaccaaggaa gtgaaagaag tggcaacgct gtctctgtgt cacaatgttt      480
40 ctggtgaaga gctggcacia actcgcatct actygcacaa ggagaagaaa atggtgctga      540
41 ctatgatgic tggggacatg aatatatggc ccgagtacaa gaaccggacc atctttgata      600
42 tcaactaata cctctccatt gtgacccctg ctctgcgcgc atctgacgag ggcacatacg      660
43 agtgtgttgt tctgaagtat gaaaaagacg ctttcaagcg ggaacacctg gctgaagtga      720
44 cgttatcagt caaagctgac ttccctacac ctagtataac tgactttgaa attccaactt      780
45 ctaaatattag aaggataatt tgcctaacct ctggagggtt tccagagcct cacctctcct      840
46 gglttgaaaa tggagaagaa ttaaatgcc acaacacaac agtttcccaa gatcctgaaa      900
47 ctgagctcta tgcctgttagc agcaaaactg atttcaatat gacaaaccaac cacagcttca      960
48 tglgtctcat caaglatgga catttaagag tgaatcagac ctccaactgg aatacaacca      1020
49 aqcaagagca ttttctgat aacctgtctc catcctgggc cattacctta atctcagtaa      1080
50 atggaaatgt tgtgatatgc tgcctgacct actgctttgc cccaagatgc agagagagaa      1140
51 ggaggaatga gagattgaga agggaaaagt tacgcccctg ataacagtgt ccgcagaagc      1200
52 aaggggctga aaagatctga aggtagcctc cglcatctct tctgggatac atggatcgtg      1260
53 gggatcatga ggcattcttc ccttaacaaa tttaagctgt tttaaccaact acctcacctt      1320
54 cttaaaaacc tctttcagat taagctgaac agttacaaga tggctggcat cctctctcct      1380
55 tctcccatac tgcaatttgc ttaatglaac ctcttctttt gccatgtttc cattctgcca      1440
56 tcttgaattg tcttgtcagc caattcatta tctattaaac actaatttga g          1491
58 <210> SEQ ID NO: 2
59 <211> LENGTH: 288
60 <212> TYPE: PRT

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61 <213> ORGANISM: Homo Sapiens
63 <400> SEQUENCE: 2
64 Met Gly His Thr Arg Arg Gln Gly Thr Ser Pro Ser Lys Cys Pro Tyr
65 1 5 10 15
66 Leu Asn Phe Phe Gln Leu Leu Val Leu Ala Gly Leu Ser His Phe Cys
67 20 25 30
68 Ser Gly Val Ile His Val Thr Lys Glu Val Lys Glu Val Ala Thr Leu
69 35 40 45
70 Ser Cys Gly His Asn Val Ser Val Glu Glu Leu Ala Gln Thr Arg Ile
71 50 55 60
72 Tyr Trp Gln Lys Glu Lys Lys Met Val Leu Thr Met Met Ser Gly Asp
73 65 70 75 80
74 Met Asn Ile Trp Pro Glu Tyr Lys Asn Arg Thr Ile Phe Asp Ile Thr
75 85 90 95
76 Asn Asn Leu Ser Ile Val Ile Leu Ala Leu Arg Pro Ser Asp Glu Gly
77 100 105 110
78 Thr Tyr Glu Cys Val Val Leu Lys Tyr Glu Lys Asp Ala Phe Lys Arg
79 115 120 125
80 Glu His Leu Ala Glu Val Thr Leu Ser Val Lys Ala Asp Phe Pro Thr
81 130 135 140
82 Pro Ser Ile Ser Asp Phe Glu Ile Pro Thr Ser Asn Ile Arg Arg Ile
83 145 150 155 160
84 Ile Cys Ser Thr Ser Gly Gly Phe Pro Glu Pro His Leu Ser Trp Leu
85 165 170 175
86 Glu Asn Gly Glu Glu Leu Asn Ala Ile Asn Thr Thr Val Ser Gln Asp
87 180 185 190
88 Pro Glu Thr Glu Leu Tyr Ala Val Ser Ser Lys Leu Asp Phe Asn Met
89 195 200 205
90 Thr Thr Asn His Ser Phe Met Cys Leu Ile Lys Tyr Gly His Leu Arg
91 210 215 220
92 Val Asn Gln Thr Phe Asn Trp Asn Thr Thr Lys Gln Glu His Phe Pro
93 225 230 235 240
94 Asp Asn Leu Leu Pro Ser Trp Ala Ile Thr Leu Ile Ser Val Asn Gly
95 245 250 255
96 Ile Phe Val Ile Cys Cys Leu Thr Tyr Cys Phe Ala Pro Arg Cys Arg
97 260 265 270
98 Glu Arg Arg Arg Asn Glu Arg Leu Arg Arg Glu Ser Val Arg Pro Val
99 275 280 285
101 <210> SEQ ID NO: 3
102 <211> LENGTH: 1424
103 <212> TYPE: DNA
104 <213> ORGANISM: Homo Sapiens
106 <400> SEQUENCE: 3
107 aggagcctta ggaggtaagg ggagctcgca aatactcctt ttggtttatt cttaccacct 60
108 tgccttctgtg ttcccttgga atgctgctgt gcttatgcat ctggtctctt tttggagcta 120
109 cagtggacag gcatttgtga cagcactatg ggactgagta acattctctt tgtgatggcc 180
110 ttccctgctct ctggtgctgc tctctgaag attcaagctt atttcaatga gactgcagac 240
111 ctgccatgcc aatttgcaaa ctctcaaac caaagcctga gtgagctagt agtattttgg 300
112 caggaccagg aaaacttggg tctgaatgag gtatacttag gcaaagagaa atttgacagt 360

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113 gttcattcca agtatatggg ccgcacaagt tttagattcgg acagttggac cctgagacttt      420
114 cacaatcttc agatcaagga caagggcttg tatcaatgta tcatccatca caaaaagccc      480
115 acaggaatga ttgcgcatcca ccagatgaat tctgaactgt cagtgccttc taacttcaqt      540
116 caacctgaaa tagtaccaat ttctaataata acagaaaatg tgtacataaa tttagacctgc      600
117 tcatctatac acggttaccc agaacctaaq aagatgagtg ttttgctaaq aaccaagaat      660
118 tcaactatcg agtatgatgg tattatgcag aaatctcaag ataattgtcac agaactgtac      720
119 gacgtttcca tcagcttgtc tgtttcattc cctgatgta cagagcaatat gaccatcttc      780
120 tgtattctgg aaactgacaa gacgcggctt ttatcttcac ctttctctat agagcttgag      840
121 gacctcagc cteccccaga ccacattcct tggattacag ctgtacttcc aacagttatt      900
122 atatgtgtga tggttttctg tctaattcta tggaaatgga agaagaagaa gcggcctcgc      960
123 aactcttata aatgtggaac caacacaatg gagaggggaag agagtgaaca gaccaagaaa      1020
124 agagaaaaaa tccatatacc tgaaagatct gatgaagccc agcgtgtttt taaaaattcg      1080
125 aagacatctt catgcgacaa aagtgtatca tgtttttaat taaagagtaa agcccataca      1140
126 agtattcatt tttctaccc ttctcttctg aagttcctgg gcaacctttt tgatttcttc      1200
127 cagaaggcaa aaagacatta ccattgagtaa laagggggct ccaggactcc ctctaagtgg      1260
128 aatagcctcc ctgttaactc agctctgctc cgtatgcaa gagagagactt taattctctt      1320
129 actgcttctt ttcacttcag agcacactta tgggccaagc ccagcttaat ggcctcatgac      1380
130 ctggaaaataa aatttaggac caataaaaaa aaaaaaaaaa aaaa      1424
132 <210> SEQ ID NO: 4
133 <211> LENGTH: 323
134 <212> TYPE: PRT
135 <213> ORGANISM: Homo Sapiens
137 <400> SEQUENCE: 4
138 Met Gly Leu Ser Asn Ile Leu Phe Val Met Ala Phe Leu Leu Ser Gly
139 1 5 10 15
140 Ala Ala Pro Leu Lys Ile Glu Ala Tyr Phe Asn Glu Thr Ala Asp Leu
141 20 25 30
142 Pro Cys Gln Phe Ala Asn Ser Gln Asn Gln Ser Leu Ser Glu Leu Val
143 35 40 45
144 Val Phe Trp Gln Asp Gln Glu Asn Leu Val Leu Asn Glu Val Tyr Leu
145 50 55 60
146 Gly Lys Glu Lys Phe Asp Ser Val His Ser Lys Tyr Met Gly Arg Thr
147 65 70 75 80
148 Ser Phe Asp Ser Asp Ser Trp Thr Leu Arg Leu His Asn Leu Gln Ile
149 85 90 95
150 Lys Asp Lys Gly Leu Tyr Gln Cys Ile Ile His His Lys Lys Pro Thr
151 100 105 110
152 Gly Met Ile Arg Ile His Gln Met Asn Ser Glu Leu Ser Val Leu Ala
153 115 120 125
154 Asn Phe Ser Gln Pro Glu Ile Val Pro Ile Ser Asn Ile Thr Glu Asn
155 130 135 140
156 Val Tyr Ile Asn Leu Thr Cys Ser Ser Ile His Gly Tyr Pro Glu Pro
157 145 150 155 160
158 Lys Lys Met Ser Val Leu Leu Arg Thr Lys Asn Ser Thr Ile Glu Tyr
159 165 170 175
160 Asp Gly Ile Met Gln Lys Ser Gln Asp Asn Val Thr Glu Leu Tyr Asp
161 180 185 190
162 Val Ser Ile Ser Leu Ser Val Ser Phe Pro Asp Val Thr Ser Asn Met
163 195 200 205

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164 Thr Ile Phe Cys Ile Leu Glu Thr Asp Lys Thr Arg Leu Leu Ser Ser
165      210      215      220
166 Pro Phe Ser Ile Glu Leu Glu Asp Pro Gln Pro Pro Pro Asp His Ile
167      225      230      235      240
168 Pro Trp Ile Thr Ala Val Leu Pro Thr Val Ile Ile Cys Val Met Val
169      245      250      255
170 Phe Cys Leu Ile Leu Trp Lys Trp Lys Lys Lys Lys Arg Pro Arg Asn
171      260      265      270
172 Ser Tyr Lys Cys Gly Thr Asn Thr Met Glu Arg Glu Glu Ser Glu Glu
173      275      280      285
174 Thr Lys Lys Arg Glu Lys Ile His Ile Pro Glu Arg Ser Asp Glu Ala
175      290      295      300
176 Gln Arg Val Phe Lys Ser Ser Lys Thr Ser Ser Cys Asp Lys Ser Asp
177      305      310      315      320
178 Thr Cys Phe
181 <210> SEQ ID NO: 5
182 <211> LENGTH: 924
183 <212> TYPE: DNA
184 <213> ORGANISM: Homo Sapiens
186 <400> SEQUENCE: 5
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188 ggaatagcgg cgtgcttgcc ggacgtgac accttccccc tggacacggc caaagtcagg      120
189 ctccaggtcc aaggtgaatg ccgacgtcc agtgttatta ggtataaagg tgtctggga      180
190 acaatcaccg ctgtggtaaa aacagaaggg cggatgaaac tctacagcgg gctgcctgcg      240
191 gggcttcagc ggcacatcag ctccgcctct ctcaagatcg gcctctacga cacgggccag      300
192 gagttcttca ccgcaggga agaaaacgca cctagttag gaagcaagat tttagctggg      360
193 ctaacgactg gaggagtggc agtattcatt gggcaacca cagaggctgt gaaagtcaga      420
194 cttcaagcac agagccatct ccacggaatc aaacctcgt acacggggac ttataatgcg      480
195 tacaqaataa tagcaacaac cgaaggcttg acgggtcttt ggaaggggac tactcccaat      540
196 ctgatgagaa gtgtcatcat caattgtaca gagctagtaa catatgatct aatgaaggag      600
197 gcttttgtga aaaacaacat attagcagat gacgtccctt gccacttggt gtrggctctt      660
198 atcgctggat tttgcgcaac agctatgtcc tccccgtgg atgtagtaaa aaccagattt      720
199 attaatcttc caccaggaca gtacaaaagt gtgcccaact gtgcaatgaa agtgttcaat      780
200 aacgaaggac caacggcttt ctccaagggg ttggtacctt ccttcttgcg acttggatec      840
201 tggaaactca ttatgtttgt gtgctttgaa caactgaaac gagaactgtc aaagtcaagg      900
202 cagactatgg actgtgccac ataa
204 <210> SEQ ID NO: 6
205 <211> LENGTH: 307
206 <212> TYPE: PRT
207 <213> ORGANISM: Homo Sapiens
209 <400> SEQUENCE: 6
210 Met Gly Gly Leu Thr Ala Ser Asp Val His Pro Thr Leu Gly Val Gln
211      1      5      10      15
212 Leu Phe Ser Ala Gly Ile Ala Ala Cys Leu Ala Asp Val Ile Thr Phe
213      20      25      30
214 Pro Leu Asp Thr Ala Lys Val Arg Leu Gln Val Gln Gly Glu Cys Pro
215      35      40      45
216 Thr Ser Ser Val Ile Arg Tyr Lys Gly Val Leu Gly Thr Ile Thr Ala
217      50      55      60

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218 Val Val Lys Thr Glu Gly Arg Met Lys Leu Tyr Ser Gly Leu Pro Ala
219 65 70 75 80
220 Gly Leu Gln Arg Gln Ile Ser Ser Ala Ser Leu Arg Ile Gly Leu Tyr
221 85 90 95
222 Asp Thr Val Gln Glu Phe Leu Thr Ala Gly Lys Glu Thr Ala Pro Ser
223 100 105 110
224 Leu Gly Ser Lys Ile Leu Ala Gly Leu Thr Thr Gly Gly Val Ala Val
225 115 120 125
226 Phe Ile Gly Gln Pro Thr Glu Val Val Lys Val Arg Leu Gln Ala Gln
227 130 135 140
228 Ser His Leu His Gly Ile Lys Pro Arg Tyr Thr Gly Thr Tyr Asn Ala
229 145 150 155 160
230 Tyr Arg Ile Ile Ala Thr Thr Glu Gly Leu Thr Gly Leu Trp Lys Gly
231 165 170 175
232 Thr Thr Pro Asn Leu Met Arg Ser Val Ile Ile Asn Cys Thr Glu Leu
233 180 185 190
234 Val Thr Tyr Asp Leu Met Lys Glu Ala Phe Val Lys Asn Asn Ile Leu
235 195 200 205
236 Ala Asp Asp Val Pro Cys His Leu Val Ser Ala Leu Ile Ala Gly Phe
237 210 215 220
238 Cys Ala Thr Ala Met Ser Ser Pro Val Asp Val Val Lys Thr Arg Phe
239 225 230 235 240
240 Ile Asn Ser Pro Pro Gly Gln Tyr Lys Ser Val Pro Asn Cys Ala Met
241 245 250 255
242 Lys Val Phe Thr Asn Glu Gly Pro Thr Ala Phe Phe Lys Gly Leu Val
243 260 265 270
244 Pro Ser Phe Leu Arg Leu Gly Ser Trp Asn Val Ile Met Phe Val Cys
245 275 280 285
246 Phe Glu Gln Leu Lys Arg Glu Leu Ser Lys Ser Arg Gln Thr Met Asp
247 290 295 300
248 Cys Ala Thr
249 305

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251 <210> SEQ ID NO: 7

252 <211> LENGTH: 1105

253 <212> TYPE: DNA

254 <213> ORGANISM: Homo Sapiens

256 <400> SEQUENCE: 7

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258 tcctgggacg tagcaggaaa tcagcatcat ggttgggttc aaggccacag atgtgcccc 120
259 tactgccact gtgaagtttc ttggggctgg cacagctgcc tgcacgcag atctcatcac 180
260 ctctctctg gatactgcta aagtcgggtt acagatccaa ggagaaagtc aggggccagt 240
261 gcgcgctaca gccagcgccc agtacgcgg tgtgatgggc accattctga ccatggtgag 300
262 tactgagggc ccccgagcc tctacaatgg gctggttgcc ggcctgcagc gccaaatgag 360
263 ctctgctct gtccgcacg gcctgtatga ttctgtcaaa cagttctaca ccaagggctc 420
264 tgagcatgcc agcattggga gccgcctcct agcaggcagc accacaggtg ccctggctgt 480
265 ggcctgtggc cagccacag atgtgtgtaa ggtccgattc caagctcagg cccgggctg 540
266 aggtggtcgg agataccaaa gcaccgtcaa tgctacaag accattgcc gagaggaag 600
267 gttccggggc ctctggaaag ggacctctc caatgtgct cgtaatgcca ttgtcaactg 660
268 tgctgagctg gtyacctatg acctcalcaa ggatgccctc ctgaaagcca acctcatgac 720

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Input Set : A:\V1397028.txt
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L:13 M:270 C: Current Application Number differs, Replaced Current Application NO
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

1

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10

24, 32 34-3460

15

~~1/11 Post 11
Jelmer
Kene~~

✓ 20

25

39. 6 - et cpi $\Delta x_{p105} \Rightarrow f_{as} \pi \Rightarrow$...

30

pg 7
tumor cells
not adhering

PMA.

pg 12 Nadi Faz Igand bawg tawar cel
melanoma, colm carummi

Fig 23 & 24
New Beach

34. The method of claim 29, wherein the apoptotic chemotherapeutic agent is from the group consisting of **adriamycin**, cytarabine, doxorubicin, and

HL 60002
Pg 70 human protein sequence alignment

[illegible]

15

20.

25

omit membrane potential

proto. motor force

A

administering an MHC class II HLA-DR ligand to the subject to selectively engage MHC class II HLA-DR on the surface of the cell in an amount effective to decrease mitochondrial membrane potential in the mammalian cell.